
Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: [year=2008; month=6; day=12; hr=19; min=27; sec=32; ms=818;]

Validated By CRFValidator v 1.0.3

Application No: 10566827 Version No: 1.1

Input Set:

Output Set:

Started: 2008-06-12 19:25:59.156

Finished: 2008-06-12 19:26:01.131

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 975 ms

Total Warnings: 20

Total Errors: 0

No. of SeqIDs Defined: 22

Actual SeqID Count: 22

Error code		Error Description									
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(1)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(2)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(3)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(4)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(5)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(6)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(7)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(8)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(9)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(10)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(11)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(12)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(13)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(14)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(17)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(18)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(19)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(20)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(21)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(22)

Input Set:

Output Set:

Started: 2008-06-12 19:25:59.156

Finished: 2008-06-12 19:26:01.131

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 975 ms

Total Warnings: 20
Total Errors: 0

No. of SeqIDs Defined: 22

Actual SeqID Count: 22

Error code Error Description

This error has occured more than 20 times, will not be displayed

SEQUENCE LISTING

```
<110> de Lorenzo Prieto, Victor
      Fernandez Herrero, Luis A
<120> System for the Production of Dimeric Proteins Based on the
       Transport System of Hemolysin of Escherichia Coli
<130> 020884-000001
<140> 10/566,827
<141> 2006-01-31
<150> P200301830 (ES)
<151> 2003-07-31
<150> PCT/ES2004/070053
<151> 2004-07-19
<160> 22
<170> PatentIn version 3.4
<210> 1
<211> 36
<212> PRT
<213> Artificial
<220>
<223> EHlyA polypeptide containing 23 kDa ('hlyA) secretion signal of
       E. coli Hly transporter tagged with the E epitope.
<400> 1
Met Thr Met Ile Thr Asn Leu Asp Leu Asn Ser Val Ser Thr Pro Gly
               5
                                   10
Gly Ala Pro Val Pro Tyr Pro Asp Pro Leu Glu Pro Ala Gly Glu Asn
                               25
            20
Ser Leu Ala Lys
       35
<210> 2
<211> 74
<212> PRT
<213> Artificial
<220>
<223> ZEHlyA polypeptide containing the 23 kDa ('hlyA) secretion signal
       of E. coli Hly transporter tagged with the E epitope.
```

<400> 2

Met Thr Met Ile Thr Asn Leu Asp Leu Asn Ser Val Ser Thr Ser Gly $1 \ 5 \ 5 \ 10 \ 15$

Gly Pro Lys Pro Ser Thr Pro Pro Gly Ser Ser Arg Met Lys Leu Glu 20 25 30

Asp Lys Val Glu Glu Leu Leu Ser Lys Asn Tyr His Leu Glu Asn Glu 35 40 45

Val Ala Arg Leu Lys Lys Leu Val Gly Glu Arg Gly Gly His His His 50 55 60

His His Ser Thr Pro Gly Gly Ala Pro 65 70

<210> 3

<211> 949

<212> DNA

<213> Artificial

<220>

<223> Ampicillin resistant plasmid pZEHlyA (sense strand); 23-kDa C-terminal domain of HlyA with E-tag epitope incorporated at the 23-kDa C domain of HlyA

<400> 3

agcggataac	aatttcacac	aggaaacagc	tatgaccatg	attacgaatt	tagatctgaa	60
tteggtgteg	acgtccggcg	gtccgaagcc	ttccactccg	cccgggtctt	cccgtatgaa	120
acagctggaa	gacaaagtag	aggagctcct	tagcaagaac	taccatctag	aaaacgaggt	180
agctcgtctg	aaaaagcttg	ttggtgaacg	tggtggtcac	catcaccatc	accatgcgtc	240
gacgcccggg	ggtgcgccgg	tgccgtatcc	ggatccgctg	gaaccggccg	gggaaaattc	300
tcttgctaaa	aatgtattat	ccggtggaaa	aggtaatgac	aagttgtacg	gcagtgaggg	360
agcagacctg	cttgatggcg	gagaagggaa	tgatcttctg	aaaggtggat	atggtaatga	420
tatttatcgt	tatctttcag	gatatggcca	tcatattatt	gacgatgaag	gggggaaaga	480
cgataaactc	agtttagctg	atatagattt	ccgggacgtt	gcctttaagc	gagaagggaa	540
tgacctcatt	atgtataaag	ctgaaggtaa	tgttctttct	attggccaca	aaaatggtat	600
tacatttaaa	aactggtttg	aaaaagagtc	agatgatctc	tctaatcatc	agatagagca	660
gatttttgat	aaagacggca	gggtaatcac	accagattct	cttaaaaaag	catttgaata	720
tcagcagagt	aataacaagg	taagttatgt	gtatggacat	gatgcatcaa	cttatgggag	780

ccaggacaat cttaatccat taattaatga aatcagcaaa atcatttcag ctgcaggtaa 840
cttcgatgtt aaggaggaaa gatctgccgc ttctttattg cagttgtccg gtaatgccag 900
tgatttttca tatggacgga actcaataac tttgacagca tcagcataa 949

<210> 4

<211> 918

<212> DNA

<213> Artificial

<220>

<223> Ampicillin resistant plasmid pZEHlyA (missense strand); 23-kDa C-terminal domain of HlyA with E-tag epitope incorporated at the 23-kDa domain of HlyA.

<400> 4

tactggtact	aatgcttaaa	tctagactta	agccacagct	gcaggccgcc	aggcttcgga	60
aggtgaggcg	ggcccagaag	ggcatacttt	gtcgaccttc	tgtttcatct	cctcgaggaa	120
tcgttcttga	tggtagatct	tttgctccat	cgagcagact	ttttcgaaca	accacttgca	180
ccaccagtgg	tagtggtagt	ggtacgcagc	tgcgggcccc	cacgcggcca	cggcataggc	240
ctaggcgacc	ttggccggcc	ccttttaaga	gaacgatttt	tacataatag	gccacctttt	300
ccattactgt	tcaacatgcc	gtcactccct	cgtctggacg	aactaccgcc	tcttccctta	360
ctagaagact	ttccacctat	accattacta	taaatagcaa	tagaaagtcc	tataccggta	420
gtataataac	tgctacttcc	cccctttctg	ctatttgagt	caaatcgact	atatctaaag	480
gccctgcaac	ggaaattcgc	tcttccctta	ctggagtaat	acatatttcg	acttccatta	540
caagaaagat	aaccggtgtt	tttaccataa	tgtaaatttt	tgaccaaact	ttttctcagt	600
ctactagaga	gattagtagt	ctatctcgtc	taaaaactat	ttctgccgtc	ccattagtgt	660
ggtctaagag	aatttttcg	taaacttata	gtcgtctcat	tattgttcca	ttcaatacac	720
atacctgtac	tacgtagttg	aataccctcg	gtcctgttag	aattaggtaa	ttaattactt	780
tagtcgtttt	agtaaagtcg	acgtccattg	aagctacaat	tcctcctttc	tagacggcga	840
agaaataacg	tcaacaggcc	attacggtca	ctaaaaagta	tacctgcctt	gagttattga	900
aactgtcgta	gtcgtatt					918

<210> 5

<211> 305

<212> PRT

<213> Artificial

<223> Ampicillin resistant plasmid pZEHlyA (protein); 23-kDa C-terminal domain of HlyA with E-tag epitope incorporated at the 23-kDa domain of HlyA.

<400> 5

Met Thr Met Ile Thr Asn Leu Asp Leu Asn Ser Val Ser Thr Ser Gly
1 5 10 15

Gly Pro Lys Pro Ser Thr Pro Pro Gly Ser Ser Arg Met Lys Gln Leu 20 25 30

Glu Asp Lys Val Glu Glu Leu Leu Ser Lys Asn Tyr His Leu Glu Asn 35 40 45

Glu Val Ala Arg Leu Lys Lys Leu Val Gly Glu Arg Gly Gly His His 50 55 60

His His His Ala Ser Thr Pro Gly Gly Ala Pro Val Pro Tyr Pro 65 70 75 80

Asp Pro Leu Glu Pro Ala Gly Glu Asn Ser Leu Ala Lys Asn Val Leu 85 90 95

Ser Gly Gly Lys Gly Asn Asp Lys Leu Tyr Gly Ser Glu Gly Ala Asp 100 105 110

Leu Leu Asp Gly Gly Glu Gly Asn Asp Leu Leu Lys Gly Gly Tyr Gly
115 120 125

Asn Asp Ile Tyr Arg Tyr Leu Ser Gly Tyr Gly His His Ile Ile Asp 130 135 140

Arg Asp Val Ala Phe Lys Arg Glu Gly Asn Asp Leu Ile Met Tyr Lys 165 170 175

Ala Glu Gly Asn Val Leu Ser Ile Gly His Lys Asn Gly Ile Thr Phe 180 185 190

Lys Asn Trp Phe Glu Lys Glu Ser Asp Asp Leu Ser Asn His Gln Ile 195 200 205 Glu Gln Ile Phe Asp Lys Asp Gly Arg Val Ile Thr Pro Asp Ser Leu 210 215 220

Lys Lys Ala Phe Glu Tyr Gln Gln Ser Asn Asn Lys Val Ser Tyr Val 225 230 235 235

Tyr Gly His Asp Ala Ser Thr Tyr Gly Ser Gln Asp Asn Leu Asn Pro 245 250 255

Leu Ile Asn Glu Ile Ser Lys Ile Ile Ser Ala Ala Gly Asn Phe Asp 260 265 270

Val Lys Glu Glu Arg Ser Ala Ala Ser Leu Leu Gln Leu Ser Gly Asn 275 280 285

Ala Ser Asp Phe Ser Tyr Gly Arg Asn Ser Ile Thr Leu Thr Ala Ser 290 295 300

Ala 305

<210> 6

<211> 1979

<212> DNA

<213> Artificial

<220>

<223> Ampicillin resistant plasmid pZEHLYA2SD (sense strand); 23-kDa C-terminal domain of HlyA with E-tag epitope incorporated at the 23-kDa domain of HlyA and polylinkerfor cloning of scFv's in frame with E-tagged 'hlyA.

<400> 6

atgaatacga atttagatct gaattcgggc ccttcgaaaa ttaatacgac tcactatagg 60 gagaccacaa cggtttccct ctagaaataa ttttgtttaa ctttaagaag gagatatatc 120 catggctage acggcctcgg gggccgcgtc gacgtccggc ggtccgaagc cttccactcc 180 gcccgggtct tcccgtatga aacagctgga agacaaagta gaggagctcc ttagcaagaa 240 300 ctaccatcta gaaaacgagg tagctcgtct gaaaaagctt gttggtgaac gtggtggtca 360 ccatcaccat caccatgcgt cgacgcccgg gggtgcgccg gtgccgtatc cggatccgct ggaaccggcc ggggaaaatt ctcttgctaa aaatgtatta tccggtggaa aaggtaatga 420 480 caagttgtac ggcagtgagg gagcagacct gcttgatggc ggagaaggga atgatcttct

gaaaggtgga tatggtaatg atatttatcg ttatctttca ggatatggcc atcatattat 540 tgacgatgaa ggggggaaag acgataaact cagtttagct gatatagatt tccgggacgt 600 tgcctttaag cgagaaggga atgacctcat tatgtataaa gctgaaggta atgttctttc 660 720 tattggccac aaaaatggta ttacatttaa aaactggttt gaaaaagagt cagatgatct ctctaatcat cagatagagc agatttttga taaagacggc agggtaatca caccagattc 780 840 tcttaaaaaa gcatttgaat atcagcagag taataacaag gtaagttatg tgtatggaca tgatgcatca acttatggga gccaggacaa tcttaatcca ttaattaatg aaatcagcaa 900 aatcatttca qctqcaqqta acttcqatqt taaqqaqqaa aqatctqccq cttctttatt 960 1020 gcagttgtcc ggtaatgcca gtgatttttc atatggacgg aactcaataa ctttgacagc atcagcataa tatattaatt taaatgatag caatcttact gggctgtgcc acataagatt 1080 gctatttttt tggagtcata atggattctt gtcataaaat tgattatggg ttatacgccc 1140 1200 tggagatttt agcccaatac cataacgtct ctgttaaccc ggaagaaatt aaacatagat 1260 ttgacacaga cgggactggt ctgggattaa cgtcatggtt gcttgctgcg aaatctttag aactaaaggt aaaacaggta aaaaaaacaa ttgaccgatt aaactttatt tctctgcccg 1320 cattagtctg gagagaggat ggacgtcatt ttattctgac taaagtcagt aaagaagcaa 1380 acagatatet tatttetgat etggageage gaaateeeeg tgttetegaa eagtetgagt 1440 ttgaggcgtt atatcagggg catattattc ttatcgcttc ccgttcttct gttgccggga 1500 aactggcgaa atttgacttt acctggttta ttcctgccat tataaaatac aggagaatat 1560 ttattgaaac ccttgttgtg tctgtttttt tacaattatt tgcattaata acccccttt 1620 tttttcaggt ggttatggac aaagtattag tgcacagggg attttcaact cttaatgtta 1680 ttactgtcgc attatctgtt gtggtggtgt ttgagattat actcagcggt ttaagaactt 1800 acatttttgc acatagtaca agtcggattg atgttgagtt gggtgccaaa ctcttccggc atttactggc gctaccgatc tcttattttg agagtcgtcg tgttggtgat actgttgcca 1860 gggtaagaga attagaccag atccgtaatt ttctgacagg acaggcatta acatctgttc 1920 tggacttatt attttcattc atattttttg cggtaatgtg gtattacagt ccaaagctt 1979

<210> 7

<211> 1979

<212> DNA

<213> Artificial

<223> Ampicillin resistant plasmid pZEHLYA2SD (missense strand); 23-kDa C-terminal domain of HlyA with E-tag epitope incorporated at the 23-kDa domain of HlyA and polylinkerfor cloning of scFv's in frame with E-tagged 'hlyA.

<400> 7 tacttatgct taaatctaga cttaagcccg ggaagctttt aattatgctg agtgatatcc 60 ctctqqtqtt qccaaaqqqa qatctttatt aaaacaaatt qaaattcttc ctctatataq 120 gtaccgatcg tgccggagcc cccggcgcag ctgcaggccg ccaggcttcg gaaggtgagg 180 cgggcccaga agggcatact ttgtcgacct tctgtttcat ctcctcgagg aatcgttctt 240 qatqqtaqat cttttqctcc atcqaqcaqa ctttttcqaa caaccacttq caccaccaqt 300 360 ggtagtggta gtggtacgca gctgcgggcc cccacgcggc cacggcatag gcctaggcga ccttggccgg ccccttttaa gagaacgatt tttacataat aggccacctt ttccattact 420 gttcaacatg ccgtcactcc ctcgtctgga cgaactaccg cctcttccct tactagaaga 480 540 ctttccacct ataccattac tataaatagc aatagaaagt cctataccgg tagtataata actgctactt cccccctttc tgctatttga gtcaaatcga ctatatctaa aggccctgca 600 acggaaattc gctcttccct tactggagta atacatattt cgacttccat tacaagaaag 660 ataaccggtg tttttaccat aatgtaaatt tttgaccaaa ctttttctca gtctactaga 720 qaqattaqta qtctatctcq tctaaaaact atttctqccq tcccattaqt qtqqtctaaq 780 agaatttttt cgtaaactta tagtcgtctc attattgttc cattcaatac acatacctgt 840 actacqtaqt tqaataccct cqqtcctqtt aqaattaqqt aattaattac tttaqtcqtt 900 ttagtaaagt cgacgtccat tgaagctaca attcctcctt tctagacggc gaagaaataa 960 cgtcaacagg ccattacggt cactaaaaag tatacctgcc ttgagttatt gaaactgtcg 1020 tagtegtatt atataattaa atttactate gttagaatga eeegacaegg tgtattetaa 1080 cgataaaaaa acctcagtat tacctaagaa cagtatttta actaataccc aatatgcggg 1140 acctctaaaa tcgggttatg gtattgcaga gacaattggg ccttctttaa tttgtatcta 1200 aactgtgtct gccctgacca gaccctaatt gcagtaccaa cgaacgacgc tttagaaatc 1260 ttgatttcca ttttgtccat tttttttgtt aactggctaa tttgaaataa agagacgggc 1320 qtaatcagac ctctctccta cctgcagtaa aataagactg atttcagtca tttcttcgtt 1380 tgtctataga ataaagacta gacctcgtcg ctttaggggc acaagagctt gtcagactca 1440 aactccgcaa tatagtcccc gtataataag aatagcgaag ggcaagaaga caacggccct 1500 1560 ttgaccgctt taaactgaaa tggaccaaat aaggacggta atattttatg tcctcttata

•	aataactttg	ggaacaacac	agacaaaaaa	atgttaataa	acgtaattat	tggggggaaa	1620
,	aaaaagtcca	ccaatacctg	tttcataatc	acgtgtcccc	taaaagttga	gaattacaat	1680
	aatgacagcg	taatagacaa	caccaccaca	aactctaata	tgagtcgcca	aattcttgaa	1740
	tgtaaaaacg	tgtatcatgt	tcagcctaac	tacaactcaa	cccacggttt	gagaaggccg	1800
	taaatgaccg	cgatggctag	agaataaaac	tctcagcagc	acaaccacta	tgacaacggt	1860
	cccattctct	taatctggtc	taggcattaa	aagactgtcc	tgtccgtaat	tgtagacaag	1920
,	acctgaataa	taaaagtaag	tataaaaaac	gccattacac	cataatgtca	ggtttcgaa	1979

<210> 8

<211> 302

<212> PRT

<213> Artificial

<220>

<223> Ampicillin resistant plasmid pZEHLYA2SD (protein); 23-kDa C-terminal domain of HlyA with E-tag epitope incorporated at the 23-kDa domain of HlyA and polylinkerfor cloning of scFv's in frame with E-tagged 'hlyA.

<400> 8

Met Ala Ser Thr Ala Ser Gly Ala Ala Ser Thr Ser Gly Gly Pro Lys

1 5 10 15

Pro Ser Thr Pro Pro Gly Ser Ser Arg Met Lys Gln Leu Glu Asp Lys 20 25 30

Val Glu Glu Leu Leu Ser Lys Asn Tyr His Leu Glu Asn Glu Val Ala 35 40 45

Arg Leu Lys Lys Leu Val Gly Glu Arg Gly Gly His His His His His 50 55

His Ala Ser Thr Pro Gly Gly Ala Pro Val Pro Tyr Pro Asp Pro Leu 65 70 75 80

Glu Pro Ala Gly Glu Asn Ser Leu Ala Lys Asn Val Leu Ser Gly Gly
85 90 95

Lys Gly Asn Asp Lys Leu Tyr Gly Ser Glu Gly Ala Asp Leu Leu Asp
100 105 110

Gly Gly Glu Gly Asn Asp Leu Leu Lys Gly Gly Tyr Gly Asn Asp Ile 115 120 125

Tyr Arg Tyr Leu Ser Gly Tyr Gly His His Ile Ile Asp Asp Glu Gly
130 135 140

Ala Phe Lys Arg Glu Gly Asn Asp Leu Ile Met Tyr Lys Ala Glu Gly
165 170 175

Asn Val Leu Ser Ile Gly His Lys Asn Gly Ile Thr Phe Lys Asn Trp 180 185 190

Phe Glu Lys Glu Ser Asp Asp Leu Ser Asn His Gln Ile Glu Gln Ile 195 200 205

Phe Asp Lys Asp Gly Arg Val Ile Thr Pro Asp Ser Leu Lys Lys Ala 210 215 220

Phe Glu Tyr Gln Gln Ser Asn Asn Lys Val Ser Tyr Val Tyr Gly His 225 230 235 240

Asp Ala Ser Thr Tyr Gly Ser Gln Asp Asn Leu Asn Pro Leu Ile Asn 245 250 255

Glu Ile Ser Lys Ile Ile Ser Ala Ala Gly Asn Phe Asp Val Lys Glu 260 265 270

Glu Arg Ser Ala Ala Ser Leu Leu Gln Leu Ser Gly Asn Ala Ser Asp 275 280 285

Phe Ser Tyr Gly Arg Asn Ser Ile Thr Leu Thr Ala Ser Ala 290 295 300

<210> 9

<211> 2792

<212> DNA

<213> Artificial

<220>

<223> Ampicillin resistant plasmid pVamyHLYA (sense strand) containing amplified DNA product encoding VHH amylase (Vamy); 23-kDa C-terminal domain of HlyA with E-tag epitope incorporated at the

<400> actataggga gaccacaacg gtttccctct agaaataatt ttgtttaact ttaagaagga 60 120 gatatatcca tggctcaggt gcagctggtg gagtcttggg gaggctcggt gcaggctggg gggtctctga gactctcctg cacagecect ggattcaect ccaatagetg ccgcatggae 180 240 tggtaccgcc aggctgcagg gaagcagcgc gagtgggtct catctattag tactgatggt cgcacaagct atgcagactc cgtgaagggc cgattcacca tctccaaaga caaagccaag 300 qacacqqtqt atctqcaaat qaacaqcctq aaacctqaqq acacqqccat ctattactqt 360 gccgtgagga cgaatgggta tcgtccgcaa tctcacgaat ttcgctactg gggcccgggg 420 acccaggtca ccgtctcctc aacggcctcg ggggccgcgt cgacgcccgg gggtgcgccg 480 gtgccgtatc cggatccgct ggaaccggcc ggggaaaatt ctcttgctaa aaatgtatta 540 tccggtggaa aaggtaatga caagttgtac ggcagtgagg gagcagacct gcttgatggc 600 660 ggagaaggga atgatcttct gaaaggtgga tatggtaatg atatttatcg ttatctttca ggatatggcc atcatattat tgacgatgaa ggggggaaag acgataaact cagtttagct 720 gatatagatt tccgggacgt tgcctttaag cgagaaggga atgacctcat tatgtataaa 780 gctgaaggta atgttctttc tattggccac aaaaatggta ttacatttaa aaactggttt 840 900 gaaaaagagt cagatgatct ctctaatcat cagatagagc agatttttga taaagacggc agggtaatca caccagattc tcttaaaaaa gcatttgaat atcagcagag taataacaag 960 gtaagttatg tgtatggaca tgatgcatca acttatggga gccaggacaa tcttaatcca 1020 ttaattaatg aaatcagcaa aatcatttca gctgcag